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TABLE 1

Target ALSV subgroups and oligonucleotide primers used in RT-PCR assay

Target Subgroup	Primer	Sequence (5'---3') ^a	Orientation	Position ^b
A	PA1	CTACAGCTGTTAGGTTCCCAGT	Forward SEQ ID No: 7	gp85 of env (5564-5586)
	PA2	GTCACCACTGTCGCCTATCCG	Reverse SEQ ID No: 8	gp85 of env (5772-5791)
A,B,C,D, E	PU1	CTRCARCTGYTAGGYTCCCAGT	Forward SEQ ID No: 15	gp85 of env 229 (5564-5585)
	PU2	GYCAYCACTGTCGCCTRTCCG	Reverse SEQ ID No: 16	gp85 of env (5772-5791)
A	PA10	GGCTTCAGGCCAAAAGGGGT	Forward SEQ ID No: 17	hr1 of gp85 232 (5642-5661)
	PA20	GTGCATTGCCACAGCGGTACTG	Reverse SEQ ID No: 18	hr2 of gp85 (5858-5879)
B	PB1	GGCTTTACCCCATACGATAG	Forward SEQ ID No: 19	hr1 of gp85 259 (5642-5661)
	PB2	ACACATCCTGACAGATGGACC A	Reverse SEQ ID No: 20	hr2 of gp85 (5861-5882)
C	PC1	TATTCGCCCCAAGGGCCAC	Forward SEQ ID No: 21	hr1 of gp85 238 (5642-5661)
	PC2	CCACGTCTCCACAGCGGTAAG T	Reverse SEQ ID No: 22	hr1 of gp85 (5858-5879)
	PD1	GGCTTCACCCCATACGGCAG	Forward SEQ ID No: 23	hr1 of gp85 258 (5642-5661)
	PD2	CCATACGTCTCACAGATAGA ATA	Reverse SEQ ID No: 24	hr1 of gp85 (5858-5882)
E	PE1	GGCTTCGCCCCACACTCCAA	Forward SEQ ID No: 25	hr1 of gp85 265 (5642-5661)
	PE2	GCACATCTCCACAGGTGTAAAT	Reverse SEQ ID No: 26	hr1 of gp85 (5867-5888)

^aR=A/G; Y=T/C. ^bPosition of primers corresponds to the numbering of the RNA genome of the Prague C strain of RSV (Schwartz et al., *Cell*, 32:853-869 1983).



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